## RT PCR PROTOCOL: BD BIOSCIENCES®

This protocol is adapted from a BDBiosciences protocol by the Gene Expression Lab. This protocol is for use with BD Bioscience's BD Advantage 2 PCR Enzyme System. For additional technical inquiries, contact Technical Service at 877-232-8995 or www.bdbiosciences.com

# BEFORE STARTING THE EXPERIMENT RT PCR PROTOCOL

Step A: Thermocycler Conditions Step B: Master Mix procedure

**TROUBLESHOOTING** 

## BEFORE STARTING THE EXPERIMENT

#### **Use PCR Hood**

For ~30 minutes prior to starting procedure, use the UV hood to decontaminate the hood.

#### RT PCR PROTOCOL

### Step A. Thermocycler Conditions

1. Enter the parameters listed in Table 1 into the thermocyler.

#### **Table 1: PCR Program**

Step	<b>Deactivation</b>		Anneal/Extension		Final Polymerization	<u>Hold:</u>
			(25 cyc	cles)*		
Temperature	95°C	95°C	60°C	72°C	72°C	4°C
Time (min.)	2:00	0:30	0:30	3:00**	7:00	<b>∞</b>

#### Step B. Master Mix Procedure

1. Prepare master mix for the number of reactions needed.

Component	<u>1rxn</u>
10X Advantage PCR Buffer(vortex)	2.5ul
50X dNTP Mix	0.5ul
Forward primer	1.0ul
Reward primer	1.0ul
50X Advantage 2 Polymerase system	0.5ul
DEPC-treated water	<u>17.5ul</u>
FINAL VOLUME	23.0ul

- Mix gently, add 23ul master mix to each well and add Add 2ul cDNA, add 24ul Master
- 3. Close lid, mix, briefly centrifuge, wait to 1000 rpm, then stop. Insert into Gene Amp 9700 and close lid, run samples.
- 4. Store at  $-20\alpha$

## TROUBLESHOOTING

#### 1. No products are observed

- Too few cycles may have been used try increasing the number of cycles, 3-5 additional cycles at a time.
- The extension time may not have been long enough if the gene is of a larger size, it may be necessary to increase the extension time.
- It is very possible that no products were observed because the gene target is difficult to amplify. It may be worthwhile to check the GC content of the gene and/or the secondary structure. Both of these are important inhibiting factors in amplification.

#### 2. Multiple products are observed

- If there are too many cycles, there may be nonspecific bands. It may be necessary to reduce the number of cycles.
- Perhaps the primers were not designed optimally. It may be necessary to make new ones.

<sup>\*25</sup> cycles for multiple-copy genes and med-high abundance of cDNA's. 30-35 cycles for low copy-number genes or rare cDNA's

<sup>\*\*</sup> If size of gene is >2kb, consider increasing length of annealing phase.